

ID HPCCPA standard; genomic RNA; VRL; 411 BP.
XX
AC D30046;
XX
SV D30046.1
XX
DT 13-MAY-1994 (Rel. 39, Created)
DT 07-JUL-2003 (Rel. 76, Last updated, Version 10)
XX
DE Hepatitis C virus (individual_isolate Td-3/93) gene for polyprotein
DE precursor, partial cds (core protein (carboxy terminus) and E1 envelope
DE protein (amino terminus half)).
XX
KW core protein; E1 envelope protein.
XX
OS Hepatitis C virus
OC Viruses; ssRNA positive-strand viruses, no DNA stage; Flaviviridae;
OC Hepacivirus.
XX
RN [1]
RP 1-411
RA Hotta H.;
RT ;
RL Submitted (28-APR-1994) to the EMBL/GenBank/DDBJ databases.
RL Hak Hotta, Kobe University School of Medicine, Department of Microbiology;
RL 7-5-1 Kusunoki-cho, Chuo-ku, Kobe, Hyogo 650, Japan
RL (Tel:078-341-7451(ex.3301), Fax:078-351-6347)
XX
RN [2]
RP 1-411
RA Hotta H.;
RT ;
RL Unpublished.
XX
RN [3]
RX MEDLINE; 95189942.
RX PUBMED; 7883898.
RA Hotta H., Handajani R., Ingelusida M., Soemarto W., Doi H., Miyajima H.,
RA Homma M.;
RT "Subtype analysis of hepatitis C virus in Indonesia on the basis of NS5b
region sequences";
RL J. Clin. Microbiol. 32(12):3049-3051(1994).
XX
DR GOA; Q81293.
DR SPTREMBL; Q81293; Q81293.
XX
CC Submitted (28-Apr-1994) to DDBJ by:
CC Hak Hotta
CC Kobe University School of Medicine
CC Department of Microbiology
CC 7-5-1 Kusunoki-cho, Chuo-ku
CC Kobe, Hyogo 650
CC Japan
CC Phone: 078-341-7451 x3301
CC Fax: 078-351-6347
XX
FH Key Location/Qualifiers
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FT /organism="Hepatitis C virus"
FT /isolate="Td-3/93"
FT CDS <1..>411

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FT /db_xref="SPTREMBL:Q81293"
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FT amino terminal half of the E1 envelope protein of hepatitis
FT C virus"
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FT /translation="LSCLLTPTAGLEYRNASGLYIVTNDCNSSSIVYEAQDIILHMPGC
FT DPCVRSGNTSRCWTPVSSTVAVGRAGAATASLRTHVDMMVGAATLCSALYVGDLCGALF
FT LVGQGFSWRHRQHWTVQDCNCISIYPGHLTGHRM"
XX
SQ Sequence 411 BP; 79 A; 120 C; 108 G; 104 T; 0 other;
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cacatgcctg gctgtgaccc ctgcgtacgc tctggcaaca catcaagggtt ctggaccct 180
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